CTATGGCTCTTAGCCAAAACCAAGCCAAGTTTTCCAAAGGATTCGTCGTGATGATTTGG MALSQNQAKFSKGFVVMIW -32 GTACTATTCATTGCTTGTGCTATAACTTCAACTGAAGCTAGTCTAACCAAATGCCAACAG 60 V L F I A C A I T S T E A S L T K C O O 5 -13 -1 +1 120 CTCCAGGCCTCGGCTAACAGTGGTCTGATAGGTACTTATGTACCACAATGCAAAGAAACG LOASANSGLIGTYVPOCKET 180 GGAGAGTTCGAAGAAAACAATGCTGGGGATCGACTGGTTACTGTTGGTGTGTGGATGAA G E F E E K Q C W G S T G Y C W C V D E 10 28 240 GATGGAAAAGAGATTCTAGGAACCAAGATCCGTGGATCTCCGGATTGCAGCCGCAGAAAA D G K E I L G T K I R G S P D C S R R K 48 A A L T L C Q M M Q A I I V N V P G W C 68 15 360 GGCCTCCATCGTGTAAAGCTGACGGCAGTTTTGACGAGGTTCAGTGCTGCGCAAGTAAT G P P S C K A D G S F D E V O C C A S N 88 420 GGAGAATGCTACTGTGTGGATAAGAAAGGAAAAGAACTTGAAGGCACAAGACAACAGGGA G E C Y C V D K K G K E L E G T R Q Q G 108 480 AGGCCAACCTGCGAAAGACACCTAAGCGAATGCGAGGAAGCTCGAATCAAGGCGCATTCA R P T C E R H L S E C E E A R I K A H S 20 128 540 AACAGTCTTCGTGTGAGATGTTCGTGCCAGAGTGTTTAGAAGATGGATCATATAACCCA N S L R V E M F V P E C L E D G S Y N P 148 600 GTACAGTGCTGGCCTAGCACAGGATACTGTTGGTGCGTCGATGAAGGAGGGGGTAAAGGTA V O C W P S T G Y C W C V D E G G V K V 168 25 660 CCAGGTTCCGATGTCAGATTTAAACGCCCCACATGCTAAGAAAAACACAGTGAACAAAGT PGSDVRFKRPTC---188 199 720 GGCTAGTTTCCAGATCGAAAATAACTACAAAGGATTAATAAAATGTTAAAATAATTTCTC

30 780 AATTCGGCTGTGATATATTTTTTCCAAGATAATTTAATCTGCATGTAGTTAACAGAAAAC

thyroglobulin domains with demonstrated CPI activity	instrated CPI activity
human invariant chain	LTKCQEEVSHIPAVHPGSFRPKC-DENGNYLPLQCYGSIGYCWCVFPNGTEVPNTRSR-GHHN-CSES
rat invariant chain (192-258)	KVLTKCQEEVSHIPDVHPGAFRPKV-DENGNYMPLQCHGSTGYCWCVFPNGTEVPHTKSR-GRHN-CSEP
chum salmon egg inh. HVPID	HVPIDGIFHLKTPCELARDAATHGPIGGFIPTC-DYNGQYTPEQCWGSTGYCWCVNSSGQKLPGTDTPPGSASNC
equistatin cDNA DOMAIN I	SLTKCQQLQASANSGLIGTYVPQC-KETGEFEEKQCWGSTGYCWCVDEDGKEILGTKIR-GSPD-CSRRK
equistatin purified domain I	SLSKCQQLQASANSGLIGAYVPQC-KETGEFEEKQCWGSTGYCWCVDEDGKEILGTKIR-GSPD-CSRRK
(protein sequence variants)	T
thyroglobulin domains with demon	thyroglobulin domains with demonstrated aspartic protease inhibitor activity (either domain II or III)

AALTLCQ--MMQAIIVNVPGWCGPPSC-KADGSFDEVQCCASNG----ECYCVDKKGKELEGTRQK-GRP-SCERHL SECEEARIKAHSNSLRVEMFVPEC-LEDGSYNPVQCWPSTG----YCWCVDEGGVKVPGSDVRFKRP-TC SPCEEARLQAHSNSLRVGMFVPQC-LEDGSYNPVQCWPSTG----YCWCVDEGGVKVPGSDVRFKRP-TC Ŀι equistatin purified domain III equistatin purified domain II (protein sequence variants) (protein sequence variants) equistatin cDNA DOMAIN III

AALTLCQ--MMQAIIVNVPGWCGPPSC-KADGSFDEVQCCASNG----ECYCVDKKGKELEGTRQQ-GRP-TCERHL

equistatin cDNA DOMAIN II

thyroglobulin domains with unknown protease inhibitor activity

human epithelial glycoprot (75-146) GSKLGRRAKPEGALQNNDGLYDPDC-DESGLFKAKQCNG-TS----MCWCVNTAGVRRTDKDTEIT----CSERVRTY EHILGAAGGADAQRPTLQGMFVPQC-DEYGHYVPTQCHHSTG----YCWCVDRDGRELEGSRTPPGMRPPCLST KCLKERQVALGGDEKVLGRFVPQC-DEKGNYEPQQFHGSTG----YSWCVNAIGEEIAGTKTPPGKIPAC bull frog saxiphilin(178-226) mouse nidogen (824-892)

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YVPQC-AEDGSFQTVQCQNDGRSCWCVGANGSEVLGSRQP-GRPVAC YLPQC-QDSGDYAPVQCDVQHVQCWCVDAEGMEVYGTRQL-GRPKRC FVPSC-TTEGSYEDVQCF-S-GECWCVNSWGKELPGSRVRDGQP-RC FVPAC-TSEGHFLPVQCFNSECYCVDAEGQAIPGTRSAIGKPKKC	Z15) QCPSLCEVLQSGVPSRRTSPGYSPACRAEDGGFSPVQCDPAQGSCWCVLGSGEEVPGTRVA-GSQPACESP KTRCQLEREHILGAAGGADAQRPTLQGMFVPQC-DEYGHYVPTQCHHSTGYCWCVDRDGRELEGSRTPPGMRPPCLSTVAP YGPCRREMEDTLNHLKFLNVLSPRGVHIPNC-DKKGFYKKKQCRPSKGRKRGFCWCVDKYGQPLPGYTTKGKEDVHCYSMQSK QKPGGLPCQNEMNRIGKLSKGKSLLGAFIPRC-NEEGYYKATQCHGSTGQCWCVDKYGNELAGSRKQ-GAV-SCEEEQET
9)	3-1215) KTRCQ -3 YGP QKPG
Thyroglobulin 1.1 (29-73) Thyroglobulin 1.2 (97-141) Thyroglobulin 1.5 (597-639) Thyroglobulin 1.6 (664-707)	Bovine thyroglobulin (1143-1215) Mouse entactin (844-923) KTRC Human IGF-binding protein-3 YG Human testican (305-381) QKP

consensus

4/13 **Figure 3**

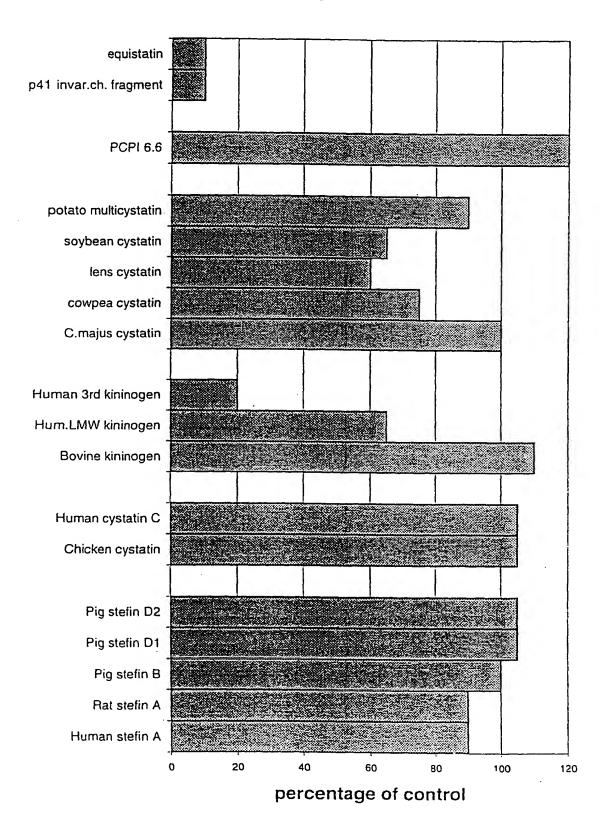


Figure 4a

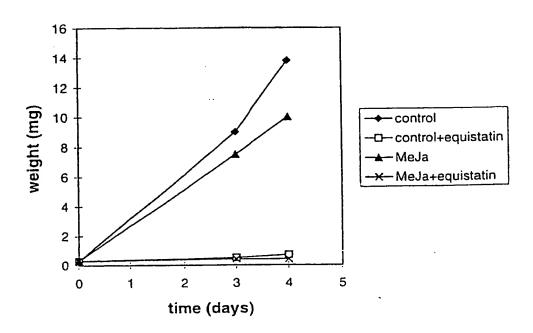
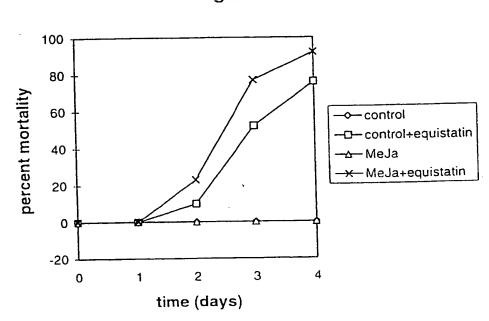
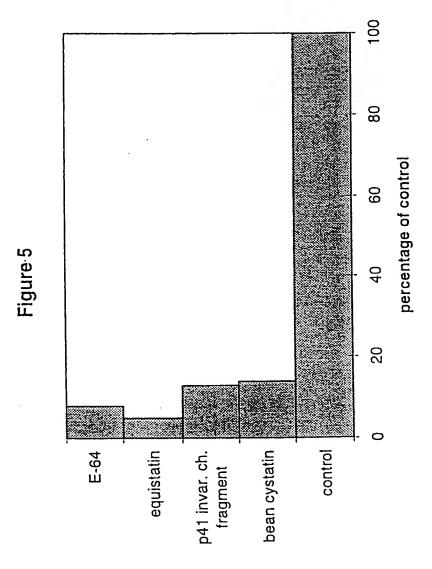
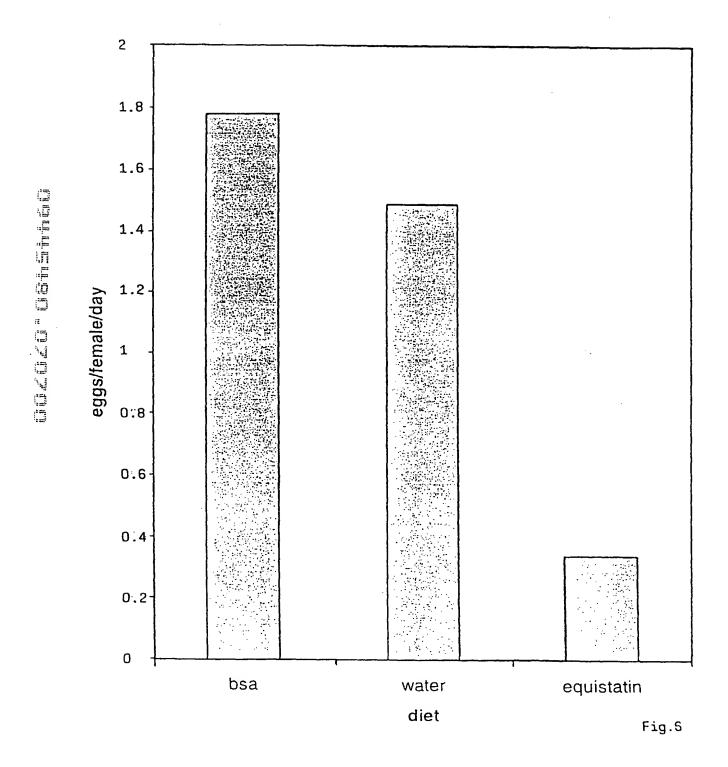


Figure 4b

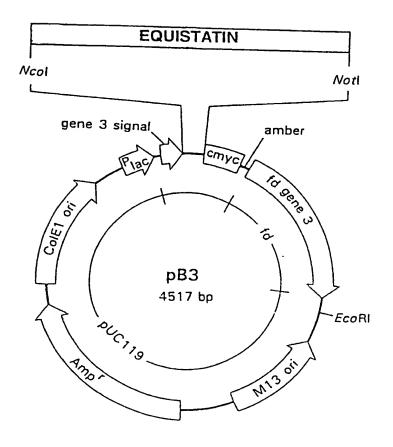




Fecundity of thrips adults on diet with equistatin (day 2)



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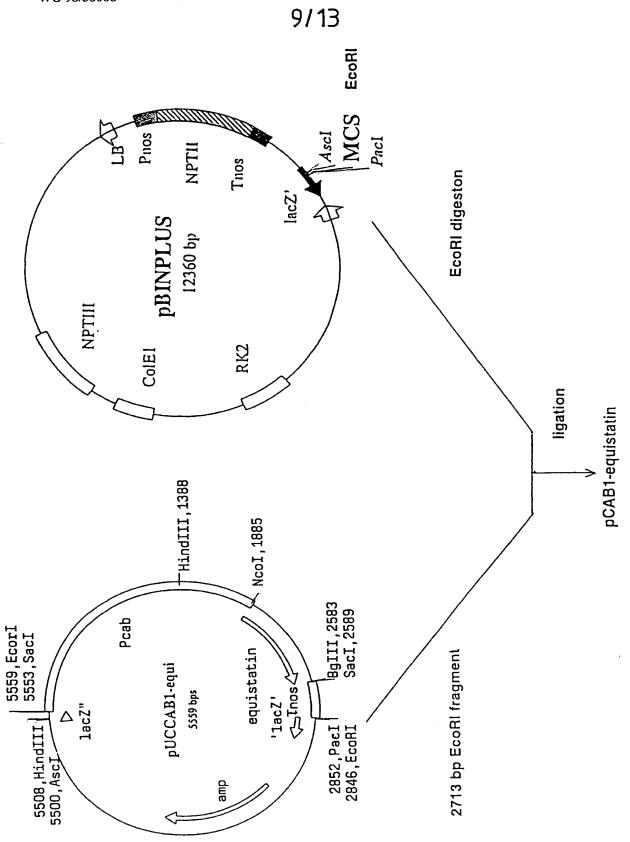


FIG. 8

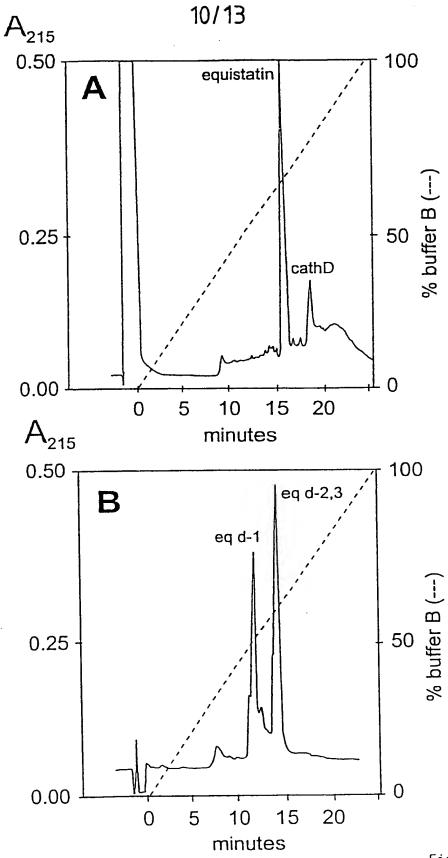


Fig.9

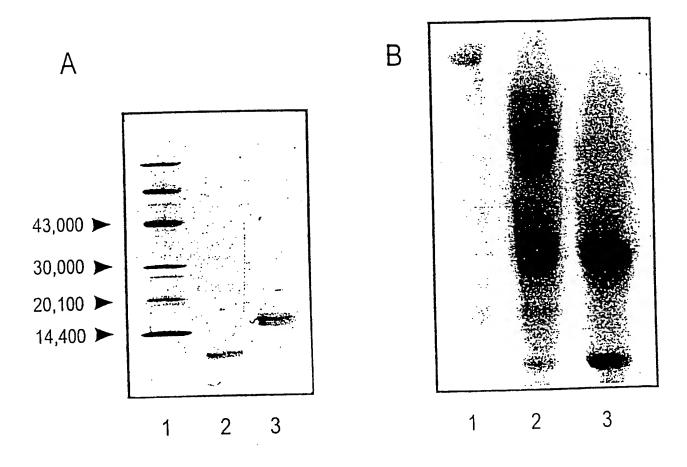
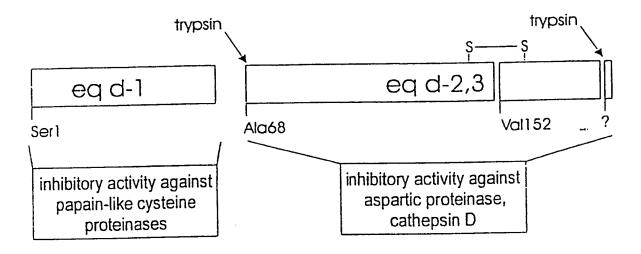


Fig.10





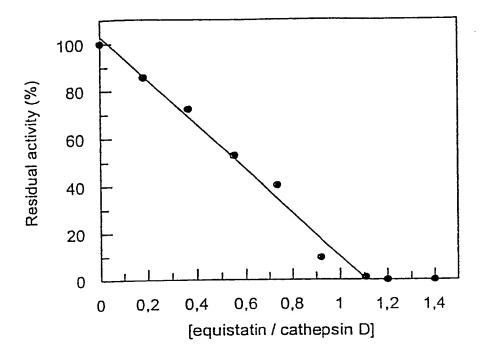


Fig.12